

SEQUENCE LISTING

<110> Welcher, Andrew A.  
Calzone, Frank J.

<120> CD20/IgE-Receptor Like Molecules and Uses Thereof

<130> 01017/36938A

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<150> US 09/723,258

<151> 2000-11-27

<150> US 60/193,728

<151> 2000-03-30

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<170> PatentIn Ver. 2.0

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<222> (98)..(697)

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Met Asp Ser Ser Thr Ala  
1 5

cac agt ccg gtg ttt ctg gta ttt cct cca gaa atc act gct tca gaa 163  
His Ser Pro Val Phe Leu Val Phe Pro Pro Glu Ile Thr Ala Ser Glu  
10 15 20

tat gag tcc aca gaa ctt tca gcc acg acc ttt tca act caa agc ccc 211  
Tyr Glu Ser Thr Glu Leu Ser Ala Thr Thr Phe Ser Thr Gln Ser Pro  
25 30 35

ttg caa aaa tta ttt gct aga aaa atg aaa atc tta ggg act atc cag 259  
Leu Gln Lys Leu Phe Ala Arg Lys Met Lys Ile Leu Gly Thr Ile Gln  
40 45 50

atc ctg ttt gga att atg acc ttt tct ttt gga gtt atc ttc ctt ttc 307  
Ile Leu Phe Gly Ile Met Thr Phe Ser Phe Gly Val Ile Phe Leu Phe  
55 60 65 70

act ttg tta aaa cca tat cca agg ttt ccc ttt ata ttt ctt tca gga 355  
Thr Leu Leu Lys Pro Tyr Pro Arg Phe Pro Phe Ile Phe Leu Ser Gly  
75 80 85

tat cca ttc tgg ggc tct gtt ttg ttc att aat tct gga gcc ttc cta 403  
Tyr Pro Phe Trp Gly Ser Val Leu Phe Ile Asn Ser Gly Ala Phe Leu  
90 95 100

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att gca gtg aaa aga aaa acc aca gaa act ctg ata ata ttg agc cga 451
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Ile Met Asn Phe Leu Ser Ala Leu Gly Ala Ile Ala Gly Ile Ile Leu
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ctc aca ttt ggt ttc atc cta gat caa aac tac att tgt ggt tat tct 547
Leu Thr Phe Gly Phe Ile Leu Asp Gln Asn Tyr Ile Cys Gly Tyr Ser
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cac caa aat agt cag tgt aag gct gtt act gtc ctg ttc ttg gga att 595
His Gln Asn Ser Gln Cys Lys Ala Val Thr Val Leu Phe Leu Gly Ile
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ttg att aca ttg atg act ttc agc att att gaa tta ttc att tct ctg 643
Leu Ile Thr Leu Met Thr Phe Ser Ile Ile Glu Leu Phe Ile Ser Leu
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cct ttc tca att ttg ggg tgc cac tca gag gat tgt gat tgt gaa caa 691
Pro Phe Ser Ile Leu Gly Cys His Ser Glu Asp Cys Asp Cys Glu Gln
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Ile Leu Gly Thr Ile Gln Ile Leu Phe Gly Ile Met Thr Phe Ser Phe
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Gly Val Ile Phe Leu Phe Thr Leu Leu Lys Pro Tyr Pro Arg Phe Pro
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Leu Ile Ile Leu Ser Arg Ile Met Asn Phe Leu Ser Ala Leu Gly Ala
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Ile Ala Gly Ile Ile Leu Leu Thr Phe Gly Phe Ile Leu Asp Gln Asn
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Gln Ser Gln Thr Met Gly Val Ser His Ser Phe Thr Pro Lys Gly Ile															
5 10 15															
act atc cct caa aga gag aaa cct gga cac atg tac caa aac gaa gat															211
Thr Ile Pro Gln Arg Glu Lys Pro Gly His Met Tyr Gln Asn Glu Asp															
20 25 30 35															
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Tyr Leu Gln Asn Gly Leu Pro Thr Glu Thr Thr Val Leu Gly Thr Val															
40 45 50															
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Gln Ile Leu Cys Cys Leu Leu Ile Ser Ser Leu Gly Ala Ile Leu Val															
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Phe Ala Pro Tyr Pro Ser His Phe Asn Pro Ala Ile Ser Thr Thr Leu															
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85 90 95															
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Ser Leu Ser Ile Ile Ser Gly Lys Gln Ser Thr Lys Pro Phe Asp Leu															
100 105 110 115															
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Ser Ser Leu Thr Ser Asn Ala Val Ser Ser Val Thr Ala Gly Ala Gly															
120 125 130															

ctc	ttc	ctc	ctt	gct	gac	agc	atg	gta	gcc	ctg	agg	act	gcc	tct	caa	547
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Ser	Asn	Asn	Pro	Gly	Ser	Ser	Phe	Ser	Ser	Thr	Gln	Ser	Gln	Asp	His	
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Ile	Gln	Gln	Val	Lys	Lys	Ser	Ser	Ser	Arg	Ser	Trp	Ile				
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Thr	Thr	Leu	Met	Ser	Gly	Tyr	Pro	Phe	Leu	Gly	Ala	Leu	Cys	Phe	Gly	

Phe Asp Leu Ser Ser Leu Thr Ser Asn Ala Val Ser Ser Val Thr Ala  
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Gly Ala Gly Leu Phe Leu Leu Ala Asp Ser Met Val Ala Leu Arg Thr  
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Ala Ser Gln His Cys Gly Ser Glu Met Asp Tyr Leu Ser Ser Leu Pro  
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Tyr Ser Glu Tyr Tyr Tyr Pro Ile Tyr Glu Ile Lys Asp Cys Leu Leu  
165 170 175

Thr Ser Val Ser Leu Thr Gly Val Leu Val Val Met Leu Ile Phe Thr  
180 185 190

Val Leu Glu Leu Leu Leu Ala Ala Tyr Ser Ser Val Phe Trp Trp Lys  
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